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TECH CENTER 1600,2000

1642

H6

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/590,583

DATE: 04/17/2001
 TIME: 11:16:27

Input Set : A:\419c9.app.txt
 Output Set: N:\CRF3\04172001\I590583.raw

ENTERED

See p.5

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3 <110> APPLICANT: Frudakis, Tony N.
4   Reed, Steven G.
5   Smith, John M.
6   Misher, Linda E.
7   Dillon, Davin C.
8   Retter, Marc W.
9   Wang, Aijun
10  Skeiky, Yasir A.W.
12 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
13   THERAPY AND DIAGNOSIS OF BREAST CANCER
16 <130> FILE REFERENCE: 210121.419C9
C--> 18 <140> CURRENT APPLICATION NUMBER: US/09/590,583
19 <141> CURRENT FILING DATE: 2000-06-08
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28 <212> TYPE: DNA
29 <213> ORGANISM: Homo sapien
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34   ttgtctaagg cgattgaagt cgtccagggg catgatgagt caccaggagt gtttttagag      180
35   cacctocagg aggcttatcg gatttacacc ccttttgacc tggcagcccc cgaaaatagc      240
36   catgctctta atttggcatt tgtggctcag gcagccccag atagtaaaag gaaactccaa      300
37   aaactagagg gattttgctg gaatgaatac cagtcagctt ttagagatag cctaaaaggt      360
38   ttt
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42 <212> TYPE: PRT
43 <213> ORGANISM: Homo sapien
45 <400> SEQUENCE: 2
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47   1          5          10          15
48   Gly Arg Thr Phe Asp Asp Phe His Arg Tyr Leu Leu Val Gly Ile Gln
49   20          25          30
50   Gly Ala Ala Gln Lys Pro Ile Asn Leu Ser Lys Ala Ile Glu Val Val
51   35          40          45
52   Gln Gly His Asp Glu Ser Pro Gly Val Phe Leu Glu His Leu Gln Glu
53   50          55          60
54   Ala Tyr Arg Ile Tyr Thr Pro Phe Asp Leu Ala Ala Pro Glu Asn Ser
55   65          70          75          80
56   His Ala Leu Asn Leu Ala Phe Val Ala Gln Ala Ala Pro Asp Ser Lys
57   85          90          95
58   Arg Lys Leu Gln Lys Leu Glu Gly Phe Cys Trp Asn Glu Tyr Gln Ser
59   100         105         110
60   Ala Phe Arg Asp Ser Leu Lys Gly Phe

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63 <210> SEQ ID NO: 3
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69 <221> NAME/KEY: misc_feature
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71 <223> OTHER INFORMATION: n = A,T,C or G
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76  tcttcaaagc ctaacagatc aagcagctct ccggtgcaca acctgcgccc aggtaaatgc      180
77  caaaaaaggt cctaaaccca gcccaggcca ccgtctccaa gaaaactcac caggagaaaa      240
78  gtgggaaatt gactttacag aagtaaaacc acaccgggct ggtacaaaat accttctagt      300
79  actggtagac accttctctg gatggactga agcatttgc accaaaaacg aaactgtcaa      360
80  tatggtagtt aagtttttac tcaatgaaat catccctcga cgtgggctgc ctgttgccat      420
81  aggtctgat  aatggaacg ccttcgcctt gtctatagtt taatcagtca gtaaggcggt      480
82  aaacattcaa tggaagctcc atttgtccta tcgaccaga gctctgggca agtagaacgc      540
83  atgaactgca ccctaaaaaa acactcttac aaaattaatc ttaaaaaccg gtgttaattg      600
84  tgtagtctc ctcccttag cctacttag agttaagggt cacccttac tgggctgggt      660
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W--> 86  gccatttgg caaaaatttc ncaactaatt tntacgtnc tacgtctccc caacagggtan      780
W--> 87  aaaaatctnc tgccttttc aaggaacat ccatccatt cctnaacaaa aggcctgccn      840
W--> 88  ttcttcccc agttaactnt ttttnttaa aattcccaa aaangaaccn cctgctggaa      900
W--> 89  aaacncccc ctccaanccc cggccnaagn ggaaggttcc ctggaatccc nccccncna      960
W--> 90  anggccgga accnttaaan tngttccngg gggtnnggcc taaaagnccn atttggtaaa      1020
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94 <211> LENGTH: 1087
95 <212> TYPE: DNA
96 <213> ORGANISM: Homo sapien
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99 <221> NAME/KEY: misc_feature
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106  ttccatcat ttaagggtt taaaatcacc ttgttcagac ctcagcatat aaaatgacct      180
107  atctgtagac ctcaggctcc aaccataccc caagagttgt ctggttttgt ttaaattact      240
108  gccaggtttc agctgcagat atccctggaa ggaatattcc agattccctg agtagtttcc      300
109  aggttaaaat cctataggct tcttctgttt tgaggaagag ttctgtcag agaaaaacat      360
110  gattttggat ttttaacttt aatgcttgtg aaacgctata aaaaaattt tctacccta      420
111  gctttaaagt actgttagtg agaaattaaa attccttcag gaggattaaa ctgccatttc      480
112  agttacccta attcctaatg ttttggtggt tagaatcttc ttaagtgtc ttgaagaagt      540
W--> 113  gttttatatt ttcccatcna gataaattct ctccnccct nttttntnt ctnntttttt      600
W--> 114  aaaacggant cttgctccgt tgtccangct gggaaatttn ttttgccaa tctccgctnc      660
W--> 115  cttgcaanaa tncgtentcc caaaattacc ncctttttcc cactccacc ccnnggaatt      720

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Input Set : A:\419c9.app.txt

Output Set: N:\CRF3\04172001\I590583.raw

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W--> 116 acctggaatt anaggccccc ncccccccc cggttaattt gtttttgttt ttagtaaaaa 780
W--> 117 acgggtttcc tgtttttagt aggatggccc anntctgacc cctnatcnt cccctcngc 840
W--> 118 cctcnaatnt tnggnntang gcttaccccc ccngnnngtt ttctctccat tnaaattttc 900
W--> 119 tntggantct tgaatnncgg gttttccctt ttaaaccnat ttttttttn nnncccccna 960
W--> 120 ttttncctcc cccntntnta angggggttt cccaanccgg gtcnccccc angtecccaa 1020
W--> 121 tttttctccc cccctctctt ttttcttnc cccaaaantc ctatcttttc cttnaaatat 1080
W--> 122 cnantnt 1087
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127 <213> ORGANISM: Homo sapien
129 <220> FEATURE:
130 <221> NAME/KEY: misc_feature
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132 <223> OTHER INFORMATION: n = A,T,C or G
134 <400> SEQUENCE: 5
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137 aggtaacaca catactatct cccaaatacc taccacaag ctcaacaatt ttaaactgtt 180
138 aggatcactg gctctaata ccatgacatg aggtcaccac caaaccatca agcgctaaac 240
139 agacagaatg ttccactcc tgatccactg tgtgggaaga agcaccgaac ttaccactg 300
W--> 140 gggggcctgc ntcanaanaa aagcccatgc ccccggtnt ncctttnaac cggaacgaat 360
W--> 141 naaccacca tccccacanc tctctgttc ntggccctg catcttggtg cctcntntnc 420
W--> 142 tttnggggan acntggggaa ggtaccccat ttcttgacc ccncnanaaa acccngtgg 480
W--> 143 cctttgccc tgattcnent gggccttttc tctttccct tttgggtgt ttaaattccc 540
W--> 144 aatgtcccn gaacctctc cntnctgcc aaaacctacc taaattntc nctangnntt 600
W--> 145 ttcttggtgt tnttttcaa aggtnacctt nctgttcan nccnacnaa aattnttcc 660
W--> 146 ntatnntggn cccnnaaaaa nnnatcnnc cnaattgcc gaattggtt gggttttcc 720
W--> 147 nctgggggaa accctttaaa ttccccctt ggcggcccc cctttttcc ccccttnga 780
W--> 148 aggcaggngg ttcttccga acttccaatt ntaacagcgn tgccattgn tgaaacctt 840
W--> 149 ttctaaaaat taaaaaatan cgggttnngg nnggcctctt tccccctcng gngggngng 900
W--> 150 aaantcctta cccnnaaaaa ggttgcttag ccccngtcc cactcccc nggaaaaatn 960
W--> 151 aaccttttcn aaaaaaggaa tataanttn cactccttn gttctctcc 1010
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155 <212> TYPE: DNA
156 <213> ORGANISM: Homo sapien
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160 <222> LOCATION: (1)...(950) /
161 <223> OTHER INFORMATION: n = A,T,C or G
163 <400> SEQUENCE: 6
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165 gctcactgca atctctgcc cgggggtcat gcgattctc tgcctcagcc ttocaagtag 120
166 ctgggattac aggcgtgcaa caccacacc ggctaatttt gtattttta tagagatgg 180
W--> 167 gttttccctt gttggccann atggtctcna accctgacc tcnngtgat ccccncccn 240
W--> 168 nganctenna ctgctgggga tnnccgnnnn nnctctccn ncncnnnnn ncncnntccn 300
W--> 169 tntctcttnc tcnnnnnnnn cnntcnntcc ncttctcnc cnnntntnt cncnncenn 360
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W--> 171 ccnntacntc ntannnnnnnt cctctntnn cctcnnnnnt cncncncnt tntctcctcn 480
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W--> 173 ctannntnnn cnnnnnnntcc ntncnttn nntcnnnnn cnnntcncn nncntnttc 600
W--> 174 ccncnnnttc cttnncntn nnnntcnnn cncntcnnn ntttctctt nntcccnnc 660
W--> 175 tcnnttcnc cnnntccnc cccncctnt ctctcncnn nntnnntnn nnnntccnc 720
W--> 176 tntcncnttc ntcnntcnc tncntcncn nncnntcnc tncntntnt ctnntcncn 780
W--> 177 tcnntntcn cctccttn ctntctctn tntctctcc ctncctnct cnttcncnc 840
W--> 178 ccnntntnn tnnncnnnt nctnnncnn cntcnnntn tctctnctn nnnntnctc 900
W--> 179 nncctncc ctntnct nctntaccn tncntcncn tctctctcc 950
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184 <213> ORGANISM: Homo sapien
186 <220> FEATURE:
187 <221> NAME/KEY: misc_feature /
188 <222> LOCATION: (1)...(1086) /
189 <223> OTHER INFORMATION: n = A,T,C or G
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194 agaaaaattc ttctgccttg agatgctgtt aatctgtaac cctagcccca accctgtgct 180
195 cacagagaca tgtgtgtgtg tgactcaagg ttcaatggat ttagggctat gctttgttaa 240
196 aaaagtgtt gaagataata tgctgttaa aagtcacac cattctctaa tctcaagtag 300
197 ccagggacac aatacactgc ggaaggccgc agggacctct gtctaggaaa gccagggtatt 360
198 gtccaagatt tctcccatg tgatagcctg agatatggcc tcatgggaag ggtaagacct 420
199 gactgtcccc cagcccgaca tccccagcc cgacatcccc cagcccgaca cccgaaaagg 480
W--> 200 gtctgtgctg aggaagatta ntaaaagagg aaggctcttt gcattgaagt aagaagaagg 540
201 ctctgtctcc tgctcgtccc tgggcaataa aatgtcttg tgttaaacc gaatgtatgt 600
202 tctacttact gagaatagga gaaaacatcc ttagggctgg aggtgagaca cctggcggc 660
W--> 203 atactgtctt ttaatgcacg agatgtttgt ntaattgcc tccagggcc nccctttcc 720
W--> 204 ttaacttttt atganacaaa aactttgttc nctttctct cgaacctct cccctattan 780
W--> 205 cctattggcc tgcccatccc ctcccaaan ggtgaaaana tgttcntaaa tncgagggaa 840
W--> 206 tccaaaacnt ttcccggtg gtccctttc caacccgct cctgggccnn ttctctccc 900
W--> 207 aacntgtccc ggntccttcn tccncncnc cttccngan aaaaaacccc gtntganggn 960
W--> 208 gccccctcaa attataacct ttcnaaaca aannngttn aaggtggtt gnttcgggtg 1020
W--> 209 cggtggcct tgagggtccc cctncacccc aatttggaan cngtttttt ttattgccc 1080
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224 aagcatcctg gagtatcaga gtttactgtt agatcagcct catttgactt cccctccac 120
225 atggtgttta aatccagcta cactacttcc tgactcaaac tccactattc ctgttcatga 180

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226 ctgtcaggaa ctgttgaaaa ctactgaaac tggccgacct gatcttcaaa atgtgcccct 240
W--> 227 aggaaagggt gatgccaccg tgttcacaga cagtaccncc ttccctcgaga agggactacg 300
W--> 228 aggggccggt gcanctgtta ccaaggagac tnatgtgttg tgggctcagg ctttaccanc 360
W--> 229 aaacacctca ncnncnaagg ctgaattgat cgcctcact caggctctcg gatggggtaa 420
230 gggatattaa cgttaacact gacagcaggt acgcctttgc tactgtgcat gtacgtggag 480
W--> 231 ccactetacca ggagcgtggg ctactcactc ggcaggtggc tgnatccac tgtaaangga 540
W--> 232 catcaaaagg aaaacnnggc tgttgcccgt ggtaaccana aanctgatcn ncagctcnaa 600
W--> 233 gatgctgtgt tgactttcac tcncncctct taaacttget gccacantc tcctttccca 660
W--> 234 accagatctg cctgacaate cccatactca aaaaaaaaaa aanactggcc ccgaaccna 720
W--> 235 accaataaaa acgggggangg tnggtnganc nncctgaccc aaaaataatg gatcccccg 780
W--> 236 gctgcaggaa ttcaattcan ccttactnat acccccacn ngngnggggg ggcngtnc 840
W--> 237 cattnccct ntattnatc tttnnccccc ccccgccnt cttttttaa ctogtgaaag 900
W--> 238 ggaaacctg ncttaccan ttatncctg gacntcccc ttccnccgtg gnttanaaaa 960
W--> 239 aaaagccnc antccntcc naaatttgca cngaaaggna aggaatttaa cctttatttt 1020
W--> 240 ttnntccttt attttgtnnn cccctttta cccagcgaa cngccatcnt ttaanaaaaa 1080
W--> 241 aaanagaang tttatttttc cttngaacca tcccaatana aancaccgc nggggaacgg 1140
W--> 242 gngngnagc cnetcacccc ctttntgtng gngggnc 1177
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251 <222> LOCATION: (1)...(1146) /
252 <223> OTHER INFORMATION: n = A,T,C or G
254 <400> SEQUENCE: 9
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256 gaaaagggtc aaaaggagct gttgacagtc atcccagggt ggccaatgtg tccagagtac 120
257 agactccatc agtgaggtca aagcctgggg cttttcagag aaggaggat tatgggtttt 180
258 ccaattatac aagtcagaag tagaaagaag ggacataaac caggaagggg gtggagcact 240
259 catcacccag agggacttgt gcctctctca gtggtagtag aggggctact tcctccacc 300
W--> 260 acggttgcaa ccaagaggca atgggtgatg agcctacagg ggacatancc gaggagacat 360
261 gggatgaccc taaggagta ggctggtttt aaggcgggtg gactgggtga gggaaactct 420
262 cctcttcttc agagagaagc agtacagggc gagctgaacc ggctgaaggt cgaggcgaaa 480
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265 gaagccggga atttcattaa caaccgcca cacagcttga acattgtgag gttcagtgac 660
W--> 266 ctttcaagg gccactccac tccaactttg gccattctac ttgcnaaat tccaaaact 720
W--> 267 tcctttttta aggccgaatc cntantccct naaaaaacnaa aaaaaatctg cncctattct 780
W--> 268 ggaaaaggcc cancccttac caggctggaa gaaattttnc cttttttttt tttttgaagg 840
W--> 269 cntttnttaa attgaacctt acccnaaaaa aaccnccng gggggcgat 900
W--> 270 ttccaaaaac naattccctt accaaaaaac aaaaaccnc ccttnttccc ttccnccctn 960
W--> 271 ttcttttaat tagggagaga tnaagcccc caatttcng gnetngatnn gtttcccccc 1020
W--> 272 ccccatctt ccnaaacttt tcccancna ggaanccnc ctttttttng gtengattna 1080
W--> 273 ncaaccttc aaacctttt tccnaaaaa ntttgnngg ngggaaaaan acctntttt 1140
W--> 274 atagan 1146
276 <210> SEQ ID NO: 10
277 <211> LENGTH: 545
278 <212> TYPE: DNA

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
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Input Set : A:\419c9.app.txt
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L:18 M:270 C: Current Application Number differs, Wrong Format
L:85 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:87 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:88 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:89 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:91 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
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L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:205 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/590,583

DATE: 04/17/2001

TIME: 11:16:28

Input Set : A:\419c9.app.txt

Output Set: N:\CRF3\04172001\I590583.raw

L:207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7

L:208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7

L:209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7

L:5725 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:322